Daniel O Stram

List of Publications by Year
in descending order

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201	24,061	14124	9346
papers	citations	69 h-index	148 g-index

204	204	204	28184
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Predicted gene expression in ancestrally diverse populations leads to discovery of susceptibility loci for lifestyle and cardiometabolic traits. American Journal of Human Genetics, 2022, 109, 669-679.	2.6	5
2	Associations of the gut microbiome with hepatic adiposity in the Multiethnic Cohort Adiposity Phenotype Study. Gut Microbes, 2021, 13, 1965463.	4.3	16
3	Ethnic Differences of Urinary Cadmium in Cigarette Smokers from the Multiethnic Cohort Study. International Journal of Environmental Research and Public Health, 2021, 18, 2669.	1.2	1
4	Genome-wide association study of pancreatic fat: The Multiethnic Cohort Adiposity Phenotype Study. PLoS ONE, 2021, 16, e0249615.	1.1	2
5	Urinary N7-(1-hydroxy-3-buten-2-yl) guanine adducts in humans: temporal stability and association with smoking. Mutagenesis, 2020, 35, 19-26.	1.0	13
6	European polygenic risk score for prediction of breast cancer shows similar performance in Asian women. Nature Communications, 2020, 11, 3833.	5.8	88
7	Minority-centric meta-analyses of blood lipid levels identify novel loci in the Population Architecture using Genomics and Epidemiology (PAGE) study. PLoS Genetics, 2020, 16, e1008684.	1.5	17
8	Genomeâ€Wide Association Study of Liver Fat: The Multiethnic Cohort Adiposity Phenotype Study. Hepatology Communications, 2020, 4, 1112-1123.	2.0	21
9	Association between mitochondrial genetic variation and breast cancer risk: The Multiethnic Cohort. PLoS ONE, 2019, 14, e0222284.	1.1	6
10	Racial/Ethnic Differences in Lung Cancer Incidence in the Multiethnic Cohort Study: An Update. Journal of the National Cancer Institute, 2019, 111, 811-819.	3.0	74
11	Genetic analyses of diverse populations improves discovery for complex traits. Nature, 2019, 570, 514-518.	13.7	679
12	Interethnic differences in pancreatic cancer incidence and risk factors: The Multiethnic Cohort. Cancer Medicine, 2019, 8, 3592-3603.	1.3	35
13	Propensity for Intra-abdominal and Hepatic Adiposity Varies Among Ethnic Groups. Gastroenterology, 2019, 156, 966-975.e10.	0.6	80
14	Breast Cancer Family History and Contralateral Breast Cancer Risk in Young Women: An Update From the Womenâ€™s Environmental Cancer and Radiation Epidemiology Study. Journal of Clinical Oncology, 2018, 36, 1513-1520.	0.8	44
15	Growth factor genes and change in mammographic density after stopping combined hormone therapy in the California Teachers Study. BMC Cancer, 2018, 18, 1072.	1.1	1
16	Evaluation of 71 Coronary Artery Disease Risk Variants in a Multiethnic Cohort. Frontiers in Cardiovascular Medicine, 2018, 5, 19.	1.1	13
17	Association of internal smoking dose with blood DNA methylation in three racial/ethnic populations. Clinical Epigenetics, 2018, 10, 110.	1.8	34
18	Tobacco biomarkers and genetic/epigenetic analysis to investigate ethnic/racial differences in lung cancer risk among smokers. Npj Precision Oncology, 2018, 2, 17.	2.3	38

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19	Estimates of Radiation Effects on Cancer Risks in the Mayak Worker, Techa River and Atomic Bomb Survivor Studies. Radiation Protection Dosimetry, 2017, 173, 26-31.	0.4	23
20	Coffee Drinking and Alcoholic and Nonalcoholic Fatty Liver Diseases and Viral Hepatitis in the Multiethnic Cohort. Clinical Gastroenterology and Hepatology, 2017, 15, 1305-1307.	2.4	22
21	BRCA2 Hypomorphic Missense Variants Confer Moderate Risks of Breast Cancer. Cancer Research, 2017, 77, 2789-2799.	0.4	75
22	Association of Common Genetic Variants With Contralateral Breast Cancer Risk in the WECARE Study. Journal of the National Cancer Institute, 2017, 109, .	3.0	28
23	Genetic Determinants of 1,3-Butadiene Metabolism and Detoxification in Three Populations of Smokers with Different Risks of Lung Cancer. Cancer Epidemiology Biomarkers and Prevention, 2017, 26, 1034-1042.	1.1	22
24	Characterizing Genetic Susceptibility to Breast Cancer in Women of African Ancestry. Cancer Epidemiology Biomarkers and Prevention, 2017, 26, 1016-1026.	1.1	24
25	Multi-SNP Haplotype Analysis Methods for Association Analysis. Methods in Molecular Biology, 2017, 1666, 485-504.	0.4	19
26	Association analysis identifies 65 new breast cancer risk loci. Nature, 2017, 551, 92-94.	13.7	1,099
27	Two Novel Susceptibility Loci for Prostate Cancer in Men of African Ancestry. Journal of the National Cancer Institute, 2017, 109, .	3.0	57
28	Impact of common genetic determinants of Hemoglobin A1c on type 2 diabetes risk and diagnosis in ancestrally diverse populations: A transethnic genome-wide meta-analysis. PLoS Medicine, 2017, 14, e1002383.	3.9	341
29	Association of CYP2A6 activity with lung cancer incidence in smokers: The multiethnic cohort study. PLoS ONE, 2017, 12, e0178435.	1.1	35
30	Metabolites of the Polycyclic Aromatic Hydrocarbon Phenanthrene in the Urine of Cigarette Smokers from Five Ethnic Groups with Differing Risks for Lung Cancer. PLoS ONE, 2016, 11, e0156203.	1.1	23
31	Prevalence of chronic liver disease and cirrhosis by underlying cause in understudied ethnic groups: The multiethnic cohort. Hepatology, 2016, 64, 1969-1977.	3.6	237
32	Breast Cancer Risk From Modifiable and Nonmodifiable Risk Factors Among White Women in the United States. JAMA Oncology, 2016, 2, 1295.	3.4	285
33	Metaâ€Analysis of Rare Variant Association Tests in Multiethnic Populations. Genetic Epidemiology, 2016, 40, 57-65.	0.6	9
34	A splicing variant of TERT identified by GWAS interacts with menopausal estrogen therapy in risk of ovarian cancer. International Journal of Cancer, 2016, 139, 2646-2654.	2.3	7
35	Genome-Wide Meta-Analyses of Breast, Ovarian, and Prostate Cancer Association Studies Identify Multiple New Susceptibility Loci Shared by at Least Two Cancer Types. Cancer Discovery, 2016, 6, 1052-1067.	7.7	157
36	Fine scale mapping of the 17q22 breast cancer locus using dense SNPs, genotyped within the Collaborative Oncological Gene-Environment Study (COGs). Scientific Reports, 2016, 6, 32512.	1.6	19

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37	Atlas of prostate cancer heritability in European and African-American men pinpoints tissue-specific regulation. Nature Communications, 2016, 7, 10979.	5.8	50
38	Prostate Cancer Susceptibility in Men of African Ancestry at 8q24. Journal of the National Cancer Institute, 2016, 108, djv431.	3.0	111
39	Genetic determinants of CYP2A6 activity across racial/ethnic groups with different risks of lung cancer and effect on their smoking intensity. Carcinogenesis, 2016, 37, 269-279.	1.3	48
40	Whole-exome sequencing of over 4100 men of African ancestry and prostate cancer risk. Human Molecular Genetics, 2016, 25, 371-381.	1.4	26
41	Breast Cancer Among Asian Americans. , 2016, , 187-218.		3
42	Benzene Uptake and Glutathione S-transferase T1 Status as Determinants of S-Phenylmercapturic Acid in Cigarette Smokers in the Multiethnic Cohort. PLoS ONE, 2016, 11, e0150641.	1.1	20
43	Lung Cancer Among Asian Americans. , 2016, , 107-136.		0
44	ABO blood group alleles and prostate cancer risk: Results from the breast and prostate cancer cohort consortium (BPC3). Prostate, 2015, 75, 1677-1681.	1.2	14
45	Fine-mapping identifies two additional breast cancer susceptibility loci at 9q31.2. Human Molecular Genetics, 2015, 24, 2966-2984.	1.4	40
46	The Contribution of Common Genetic Variation to Nicotine and Cotinine Glucuronidation in Multiple Ethnic/Racial Populations. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 119-127.	1.1	47
47	Generalizability of established prostate cancer risk variants in men of <scp>A</scp>frican ancestry. International Journal of Cancer, 2015, 136, 1210-1217.	2.3	62
48	Fine-Scale Mapping of the 5q11.2 Breast Cancer Locus Reveals at Least Three Independent Risk Variants Regulating MAP3K1. American Journal of Human Genetics, 2015, 96, 5-20.	2.6	76
49	Genome-wide association analysis of more than 120,000 individuals identifies 15 new susceptibility loci for breast cancer. Nature Genetics, 2015, 47, 373-380.	9.4	513
50	Genome-wide Analysis Identifies Novel Loci Associated with Ovarian Cancer Outcomes: Findings from the Ovarian Cancer Association Consortium. Clinical Cancer Research, 2015, 21, 5264-5276.	3.2	33
51	Integration of multiethnic fine-mapping and genomic annotation to prioritize candidate functional SNPs at prostate cancer susceptibility regions. Human Molecular Genetics, 2015, 24, 5603-5618.	1.4	50
52	Evaluating the ovarian cancer gonadotropin hypothesis: A candidate gene study. Gynecologic Oncology, 2015, 136, 542-548.	0.6	15
53	Variation in Levels of the Lung Carcinogen NNAL and Its Glucuronides in the Urine of Cigarette Smokers from Five Ethnic Groups with Differing Risks for Lung Cancer. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 561-569.	1.1	39
54	Associations Between Genetic Ancestries and Nicotine Metabolism Biomarkers in the Multiethnic Cohort Study. American Journal of Epidemiology, 2015, 182, 945-951.	1.6	12

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55	Population Distribution of Lifetime Risk of Ovarian Cancer in the United States. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 671-676.	1.1	82
56	Identification and characterization of novel associations in the CASP8/ALS2CR12 region on chromosome 2 with breast cancer risk. Human Molecular Genetics, 2015, 24, 285-298.	1.4	38
57	Mercapturic Acids Derived from the Toxicants Acrolein and Crotonaldehyde in the Urine of Cigarette Smokers from Five Ethnic Groups with Differing Risks for Lung Cancer. PLoS ONE, 2015, 10, e0124841.	1.1	56
58	Pleiotropy of Cancer Susceptibility Variants on the Risk of Non-Hodgkin Lymphoma: The PAGE Consortium. PLoS ONE, 2014, 9, e89791.	1.1	16
59	Multiple Nonglycemic Genomic Loci Are Newly Associated With Blood Level of Glycated Hemoglobin in East Asians. Diabetes, 2014, 63, 2551-2562.	0.3	61
60	Nicotine N-glucuronidation relative to N-oxidation and C-oxidation and UGT2B10 genotype in five ethnic/racial groups. Carcinogenesis, 2014, 35, 2526-2533.	1.3	124
61	Additive Interactions Between Susceptibility Single-Nucleotide Polymorphisms Identified in Genome-Wide Association Studies and Breast Cancer Risk Factors in the Breast and Prostate Cancer Cohort Consortium. American Journal of Epidemiology, 2014, 180, 1018-1027.	1.6	36
62	Imputation and subset-based association analysis across different cancer types identifies multiple independent risk loci in the TERT-CLPTM1L region on chromosome 5p15.33. Human Molecular Genetics, 2014, 23, 6616-6633.	1.4	90
63	Common non-synonymous SNPs associated with breast cancer susceptibility: findings from the Breast Cancer Association Consortium. Human Molecular Genetics, 2014, 23, 6096-6111.	1.4	53
64	Trans-ethnic genome-wide association study of colorectal cancer identifies a new susceptibility locus in VTI1A. Nature Communications, 2014, 5, 4613.	5.8	72
65	The Role of Local Ancestry Adjustment in Association Studies Using Admixed Populations. Genetic Epidemiology, 2014, 38, 502-515.	0.6	38
66	Post-GWAS Analyses. Statistics in the Health Sciences, 2014, , 285-327.	0.2	0
67	A comprehensive examination of breast cancer risk loci in African American women. Human Molecular Genetics, 2014, 23, 5518-5526.	1.4	42
68	Diabetes and Racial/Ethnic Differences in Hepatocellular Carcinoma Risk: The Multiethnic Cohort. Journal of the National Cancer Institute, 2014, 106, dju326-dju326.	3.0	44
69	Design, Analysis, and Interpretation of Genome-Wide Association Scans. Statistics in the Health Sciences, 2014, , .	0.2	16
70	The Impact of GWAS Findings on Cancer Etiology and Prevention. Current Epidemiology Reports, 2014, 1, 130-137.	1.1	3
71	A meta-analysis of 87,040 individuals identifies 23 new susceptibility loci for prostate cancer. Nature Genetics, 2014, 46, 1103-1109.	9.4	408
72	Evidence that breast cancer risk at the 2q35 locus is mediated through IGFBP5 regulation. Nature Communications, 2014, 5, 4999.	5.8	105

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73	Fine-Mapping IGF1 and Prostate Cancer Risk in African Americans: The Multiethnic Cohort Study. Cancer Epidemiology Biomarkers and Prevention, 2014, 23, 1928-1932.	1.1	7
74	Joint Effects of Known Type 2 Diabetes Susceptibility Loci in Genome-Wide Association Study of Singapore Chinese: The Singapore Chinese Health Study. PLoS ONE, 2014, 9, e87762.	1.1	15
75	SNP Imputation for Association Studies. Statistics in the Health Sciences, 2014, , 213-242.	0.2	0
76	Correcting for Hidden Population Structure in Single Marker Association Testing and Estimation. Statistics in the Health Sciences, 2014, , 135-181.	0.2	1
77	An Introduction to Association Analysis. Statistics in the Health Sciences, 2014, , 79-133.	0.2	0
78	Haplotype Imputation for Association Analysis. Statistics in the Health Sciences, 2014, , 183-211.	0.2	0
79	Fine-Scale Mapping of the FGFR2 Breast Cancer Risk Locus: Putative Functional Variants Differentially Bind FOXA1 and E2F1. American Journal of Human Genetics, 2013, 93, 1046-1060.	2.6	98
80	Multiple independent variants at the TERT locus are associated with telomere length and risks of breast and ovarian cancer. Nature Genetics, 2013, 45, 371-384.	9.4	493
81	A genome-wide association study of breast cancer in women of African ancestry. Human Genetics, 2013, 132, 39-48.	1.8	70
82	Functional Variants at the 11q13 Risk Locus for Breast Cancer Regulate Cyclin D1 Expression through Long-Range Enhancers. American Journal of Human Genetics, 2013, 92, 489-503.	2.6	201
83	Genome-wide association studies identify four ER negativeâ€“specific breast cancer risk loci. Nature Genetics, 2013, 45, 392-398.	9.4	374
84	Large-scale genotyping identifies 41 new loci associated with breast cancer risk. Nature Genetics, 2013, 45, 353-361.	9.4	960
85	Common genetic determinants of breast-cancer risk in East Asian women: a collaborative study of 23 637 breast cancer cases and 25 579 controls. Human Molecular Genetics, 2013, 22, 2539-2550.	1.4	86
86	Levels of Beta-Microseminoprotein in Blood and Risk of Prostate Cancer in Multiple Populations. Journal of the National Cancer Institute, 2013, 105, 237-243.	3.0	42
87	Combined and Interactive Effects of Environmental and GWAS-Identified Risk Factors in Ovarian Cancer. Cancer Epidemiology Biomarkers and Prevention, 2013, 22, 880-890.	1.1	54
88	Dietary patterns and breast cancer risk in the California Teachers Study cohort. American Journal of Clinical Nutrition, 2013, 98, 1524-1532.	2.2	100
89	Epigenetic analysis leads to identification of HNF1B as a subtype-specific susceptibility gene for ovarian cancer. Nature Communications, 2013, 4, 1628.	5.8	144
90	A Genome-Wide Scan for Breast Cancer Risk Haplotypes among African American Women. PLoS ONE, 2013, 8, e57298.	1.1	20

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91	Association of Type 2 Diabetes Susceptibility Variants With Advanced Prostate Cancer Risk in the Breast and Prostate Cancer Cohort Consortium. American Journal of Epidemiology, 2012, 176, 1121-1129.	1.6	67
92	Prediction of breast cancer risk by genetic risk factors, overall and by hormone receptor status. Journal of Medical Genetics, 2012, 49, 601-608.	1.5	58
93	Genetic Variation in Peroxisome Proliferatorâ€“Activated Receptor Gamma, Soy, and Mammographic Density in Singapore Chinese Women. Cancer Epidemiology Biomarkers and Prevention, 2012, 21, 635-644.	1.1	16
94	Evaluating Genetic Risk for Prostate Cancer among Japanese and Latinos. Cancer Epidemiology Biomarkers and Prevention, 2012, 21, 2048-2058.	1.1	51
95	Multi-SNP Haplotype Analysis Methods for Association Analysis. Methods in Molecular Biology, 2012, 850, 423-452.	0.4	17
96	A meta-analysis of genome-wide association studies of breast cancer identifies two novel susceptibility loci at 6q14 and 20q11. Human Molecular Genetics, 2012, 21, 5373-5384.	1.4	168
97	Interactions Between Genetic Variants and Breast Cancer Risk Factors in the Breast and Prostate Cancer Cohort Consortium. Journal of the National Cancer Institute, 2011, 103, 1252-1263.	3.0	147
98	A common variant at the TERT-CLPTM1L locus is associated with estrogen receptorâ€“negative breast cancer. Nature Genetics, 2011, 43, 1210-1214.	9.4	279
99	Generalizability and Epidemiologic Characterization of Eleven Colorectal Cancer GWAS Hits in Multiple Populations. Cancer Epidemiology Biomarkers and Prevention, 2011, 20, 70-81.	1.1	73
100	Genome-wide association study of prostate cancer in men of African ancestry identifies a susceptibility locus at 17q21. Nature Genetics, 2011, 43, 570-573.	9.4	198
101	Caution in generalizing known genetic risk markers for breast cancer across all ethnic/racial populations. European Journal of Human Genetics, 2011, 19, 243-245.	1.4	17
102	Genetic variation in insulin-like growth factor 2 may play a role in ovarian cancer risk. Human Molecular Genetics, 2011, 20, 2263-2272.	1.4	22
103	Genome-wide association study identifies new prostate cancer susceptibility loci. Human Molecular Genetics, 2011, 20, 3867-3875.	1.4	160
104	No Association of Type 2 Diabetes Risk Variants and Prostate Cancer Risk: the Multiethnic Cohort and PAGE: Table 1.. Cancer Epidemiology Biomarkers and Prevention, 2011, 20, 1979-1981.	1.1	11
105	Prostate Cancer Susceptibility Polymorphism rs2660753 Is Not Associated with Invasive Ovarian Cancer. Cancer Epidemiology Biomarkers and Prevention, 2011, 20, 1028-1031.	1.1	0
106	The Role of KRAS rs61764370 in Invasive Epithelial Ovarian Cancer: Implications for Clinical Testing. Clinical Cancer Research, 2011, 17, 3742-3750.	3.2	47
107	Fine-mapping of breast cancer susceptibility loci characterizes genetic risk in African Americans. Human Molecular Genetics, 2011, 20, 4491-4503.	1.4	61
108	Characterizing Genetic Risk at Known Prostate Cancer Susceptibility Loci in African Americans. PLoS Genetics, 2011, 7, e1001387.	1.5	117

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109	Characterizing Associations and SNP-Environment Interactions for GWAS-Identified Prostate Cancer Risk Markersâ€"Results from BPC3. PLoS ONE, 2011, 6, e17142.	1.1	57
110	Self-reported ethnicity, genetic structure and the impact of population stratification in a multiethnic study. Human Genetics, 2010, 128, 165-177.	1.8	43
111	Common variants at 19p13 are associated with susceptibility to ovarian cancer. Nature Genetics, 2010, 42, 880-884.	9.4	235
112	A genome-wide association study identifies susceptibility loci for ovarian cancer at 2q31 and 8q24. Nature Genetics, 2010, 42, 874-879.	9.4	321
113	Eighteen Insulin-like Growth Factor Pathway Genes, Circulating Levels of IGF-I and Its Binding Protein, and Risk of Prostate and Breast Cancer. Cancer Epidemiology Biomarkers and Prevention, 2010, 19, 2877-2887.	1.1	59
114	Pooled Analysis of Phosphatidylinositol 3-Kinase Pathway Variants and Risk of Prostate Cancer. Cancer Research, 2010, 70, 2389-2396.	0.4	43
115	Comprehensive analysis of common genetic variation in 61 genes related to steroid hormone and insulin-like growth factor-I metabolism and breast cancer risk in the NCI breast and prostate cancer cohort consortiumâ€. Human Molecular Genetics, 2010, 19, 3873-3884.	1.4	45
116	A Common Prostate Cancer Risk Variant 5â€² of Microseminoprotein-Î² (MSMB) Is a Strong Predictor of Circulating Î²-Microseminoprotein (MSP) Levels in Multiple Populations. Cancer Epidemiology Biomarkers and Prevention, 2010, 19, 2639-2646.	1.1	17
117	A comprehensive analysis of common IGF1, IGFBP1 and IGFBP3 genetic variation with prospective IGF-I and IGFBP-3 blood levels and prostate cancer risk among Caucasians â€. Human Molecular Genetics, 2010, 19, 3089-3101.	1.4	47
118	Evaluation of Candidate Stromal Epithelial Cross-Talk Genes Identifies Association between Risk of Serous Ovarian Cancer and TERT, a Cancer Susceptibility â€œHot-Spotâ€• PLoS Genetics, 2010, 6, e1001016.	1.5	48
119	Consistent Association of Type 2 Diabetes Risk Variants Found in Europeans in Diverse Racial and Ethnic Groups. PLoS Genetics, 2010, 6, e1001078.	1.5	168
120	Exploring genetic susceptibility to cancer in diverse populations. Current Opinion in Genetics and Development, 2010, 20, 330-335.	1.5	28
121	Recent breast cancer incidence trends according to hormone therapy use: the California Teachers Study cohort. Breast Cancer Research, 2010, 12, R4.	2.2	39
122	Methodological Issues in Multistage Genome-Wide Association Studies. Statistical Science, 2009, 24, 414-429.	1.6	41
123	Generalizability of Associations from Prostate Cancer Genome-Wide Association Studies in Multiple Populations. Cancer Epidemiology Biomarkers and Prevention, 2009, 18, 1285-1289.	1.1	102
124	Association between invasive ovarian cancer susceptibility and 11 best candidate SNPs from breast cancer genome-wide association study. Human Molecular Genetics, 2009, 18, 2297-2304.	1.4	42
125	Association of Diabetes With Prostate Cancer Risk in the Multiethnic Cohort. American Journal of Epidemiology, 2009, 169, 937-945.	1.6	136
126	IGF2R Missense Single-Nucleotide Polymorphisms and Breast Cancer Risk: The Multiethnic Cohort Study. Cancer Epidemiology Biomarkers and Prevention, 2009, 18, 1922-1924.	1.1	10

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127	Quantitative trait loci predicting circulating sex steroid hormones in men from the NCI-Breast and Prostate Cancer Cohort Consortium (BPC3). Human Molecular Genetics, 2009, 18, 3749-3757.	1.4	37
128	CYP19A1 Genetic Variation in Relation to Prostate Cancer Risk and Circulating Sex Hormone Concentrations in Men from the Breast and Prostate Cancer Cohort Consortium. Cancer Epidemiology Biomarkers and Prevention, 2009, 18, 2734-2744.	1.1	33
129	Genetic polymorphisms of the GNRH1 and GNRHR genes and risk of breast cancer in the National Cancer Institute Breast and Prostate Cancer Cohort Consortium (BPC3). BMC Cancer, 2009, 9, 257.	1.1	5
130	A genome-wide association study identifies a new ovarian cancer susceptibility locus on 9p22.2. Nature Genetics, 2009, 41, 996-1000.	9.4	276
131	Dietary assessment in the California Teachers Study: reproducibility and validity. Cancer Causes and Control, 2008, 19, 595-603.	0.8	55
132	Haplotypes of the estrogen receptor beta gene and breast cancer risk. International Journal of Cancer, 2008, 122, 387-392.	2.3	38
133	Utilizing HapMap and Tagging SNPs. Methods in Molecular Medicine, 2008, 141, 37-54.	0.8	17
134	Comprehensive association testing of common genetic variation in DNA repair pathway genes in relationship with breast cancer risk in multiple populations. Human Molecular Genetics, 2008, 17, 825-834.	1.4	42
135	Dietary Patterns and Risk of Ovarian Cancer in the California Teachers Study Cohort. Nutrition and Cancer, 2008, 60, 285-291.	0.9	27
136	Heterogeneity of Breast Cancer Associations with Five Susceptibility Loci by Clinical and Pathological Characteristics. PLoS Genetics, 2008, 4, e1000054.	1.5	315
137	IGF-1, IGFBP-1, and IGFBP-3 Polymorphisms Predict Circulating IGF Levels but Not Breast Cancer Risk: Findings from the Breast and Prostate Cancer Cohort Consortium (BPC3). PLoS ONE, 2008, 3, e2578.	1.1	106
138	Sequence Variants of Estrogen Receptor Î² and Risk of Prostate Cancer in the National Cancer Institute Breast and Prostate Cancer Cohort Consortium. Cancer Epidemiology Biomarkers and Prevention, 2007, 16, 1973-1981.	1.1	33
139	Exploiting Gene-Environment Interaction to Detect Genetic Associations. Human Heredity, 2007, 63, 111-119.	0.4	387
140	Beta-Cryptoxanthin and Lung Cancer in Shanghai, Chinaâ€"An Examination of Potential Confounding with Cigarette Smoking Using Urinary Cotinine as a Biomarker for True Tobacco Exposure. Nutrition and Cancer, 2007, 57, 123-129.	0.9	7
141	Genetic Variation at the CYP19A1 Locus Predicts Circulating Estrogen Levels but not Breast Cancer Risk in Postmenopausal Women. Cancer Research, 2007, 67, 1893-1897.	0.4	140
142	Risk Factors for Renal Cell Cancer: The Multiethnic Cohort. American Journal of Epidemiology, 2007, 166, 932-940.	1.6	175
143	Re: The Use of Inferred Haplotypes in Downstream Analysis. American Journal of Human Genetics, 2007, 81, 863-865.	2.6	16
144	CYP17 Genetic Variation and Risk of Breast and Prostate Cancer from the National Cancer Institute Breast and Prostate Cancer Cohort Consortium (BPC3). Cancer Epidemiology Biomarkers and Prevention, 2007, 16, 2237-2246.	1.1	54

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145	A comprehensive analysis of common genetic variation in prolactin (PRL) and PRL receptor (PRLR) genes in relation to plasma prolactin levels and breast cancer risk: the Multiethnic Cohort. BMC Medical Genetics, 2007, 8, 72.	2.1	40
146	Multiple regions within 8q24 independently affect risk for prostate cancer. Nature Genetics, 2007, 39, 638-644.	9.4	621
147	A common genetic risk factor for colorectal and prostate cancer. Nature Genetics, 2007, 39, 954-956.	9.4	336
148	A comprehensive analysis of the androgen receptor gene and risk of breast cancer: results from the National Cancer Institute Breast and Prostate Cancer Cohort Consortium (BPC3). Breast Cancer Research, 2006, 8, R54.	2.2	32
149	Transferability of tag SNPs in genetic association studies in multiple populations. Nature Genetics, 2006, 38, 1298-1303.	9.4	224
150	An utter refutation of the â€˜Fundamental Theorem of the HapMapâ€™ by Terwilliger and Hiekkalinna. European Journal of Human Genetics, 2006, 14, 1238-1239.	1.4	6
151	Optimal two-stage genotyping designs for genome-wide association scans. Genetic Epidemiology, 2006, 30, 356-368.	0.6	119
152	Ethnic and Racial Differences in the Smoking-Related Risk of Lung Cancer. New England Journal of Medicine, 2006, 354, 333-342.	13.9	668
153	Haplotype Analysis of the HSD17B1 Gene and Risk of Breast Cancer: A Comprehensive Approach to Multicenter Analyses of Prospective Cohort Studies. Cancer Research, 2006, 66, 2468-2475.	0.4	64
154	Haplotype-Based Association Studies of IGFBP1 and IGFBP3 with Prostate and Breast Cancer Risk: The Multiethnic Cohort. Cancer Epidemiology Biomarkers and Prevention, 2006, 15, 1993-1997.	1.1	47
155	IGF-I Genetic Variation and Breast Cancer: the Multiethnic Cohort. Cancer Epidemiology Biomarkers and Prevention, 2006, 15, 172-174.	1.1	21
156	Common Genetic Variation in IGF1 and Prostate Cancer Risk in the Multiethnic Cohort. Journal of the National Cancer Institute, 2006, 98, 123-134.	3.0	107
157	Common Genetic Variation at PTEN and Risk of Sporadic Breast and Prostate Cancer. Cancer Epidemiology Biomarkers and Prevention, 2006, 15, 1021-1025.	1.1	27
158	A Systematic Assessment of Common Genetic Variation in CYP11A and Risk of Breast Cancer. Cancer Research, 2006, 66, 12019-12025.	0.4	19
159	Software for tag single nucleotide polymorphism selection. Human Genomics, 2005, 2, 144.	1.4	35
160	A controlled 2-mo dietary fat reduction and soy food supplementation study in postmenopausal women. American Journal of Clinical Nutrition, 2005, 81, 1133-1141.	2.2	47
161	Finding and using haplotype blocks in candidate gene association studies. , 2005, , .		0
162	Asymptotic equivalence between two score tests for haplotype-specific risk in general linear models. Genetic Epidemiology, 2005, 29, 166-170.	0.6	18

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163	Enlarged and prominent nucleoli may be indicative ofMYCN amplification. Cancer, 2005, 103, 174-180.	2.0	38
164	Genetic Variation in the HSD17B1 Gene and Risk of Prostate Cancer. PLoS Genetics, 2005, 1, e68.	1.5	66
165	Genetic variation in the HSD17B1 gene and risk of prostate cancer. PLoS Genetics, 2005, preprint, e68.	1.5	6
166	Clarifying the PROGINS Allele Association in Ovarian and Breast Cancer Risk: A Haplotype-Based Analysis. Journal of the National Cancer Institute, 2005, 97, 51-59.	3.0	62
167	Comparison of Prostate-Specific Antigen and Hormone Levels among Men in Singapore and the United States. Cancer Epidemiology Biomarkers and Prevention, 2005, 14, 1692-1696.	1.1	27
168	A Haplotype-Based Case-Control Study of BRCA1 and Sporadic Breast Cancer Risk. Cancer Research, 2005, 65, 7516-7522.	0.4	53
169	Common variation in BRCA2 and breast cancer risk: a haplotype-based analysis in the Multiethnic Cohort. Human Molecular Genetics, 2004, 13, 2431-2441.	1.4	51
170	Tag SNP selection for association studies. Genetic Epidemiology, 2004, 27, 365-374.	0.6	165
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